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SEQUENCE LISTING

<110> YEH, EDWARD T.H.  
<120> COMPOSITIONS AND USES FOR A NOVEL CELL-DEATH-PROTECTING  
PROTEIN

<130> UTSH:248  
<140> 09/484,964

<150> 08/964,162  
<151> 1997-11-04

<150> 60/030,302  
<151> 1996-11-05

<160> 16  
<170> PatentIn Ver. 2.0

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<212> DNA  
<213> Homo sapiens  
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Met Ser Asp Gln Glu Ala Lys Pro Ser  
1 5  
act gag gac ttg ggg gat aag aag caa ggt gaa tat att aaa ctc aaa 162  
Thr Glu Asp Leu Gly Asp Lys Lys Gln Gly Glu Tyr Ile Lys Leu Lys 25  
10 15 20  
gtc att gga cag gat agc agt gag att cac ttc aaa gtg aaa atg aca 210  
Val Ile Gly Gln Asp Ser Ser Glu Ile His Phe Lys Val Lys Met Thr 40  
30 35 40  
aca cat ctc aag aaa ctc aaa gaa tca tac tgt caa aga cag ggt gtt 258  
Thr His Leu Lys Lys Leu Lys Glu Ser Tyr Cys Gln Arg Gln Gly Val 55  
45 50 55  
cca atg aat tca ctc agg ttt ctc ttt gag ggt cag aga att gct gat 306  
Pro Met Asn Ser Leu Arg Phe Leu Phe Glu Gly Gln Arg Ile Ala Asp 70  
60 65 70  
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75 80 85  
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<211> 101  
<212> PRT  
<213> Homo sapiens

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20 25 30  
Glu Ile His Phe Lys Val Lys Met Thr Thr His Leu Lys Lys Leu Lys  
35 40 45  
Glu Ser Tyr Cys Gln Arg Gln Gly Val Pro Met Asn Ser Leu Arg Phe  
50 55 60  
Leu Phe Glu Gly Gln Arg Ile Ala Asp Asn His Thr Pro Lys Glu Leu  
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Gly Met Glu Glu Glu Asp Val Ile Glu Val Tyr Gln Glu Gln Thr Gly  
85 90 95  
Gly His Ser Thr Val  
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<210> 3  
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<222> (689)  
<223> N = A, C, G or T  
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 cagtttaaga ttaagaggca tacaccactt agtaaaactaa tgaaagccta ttgtgaacga 240  
 cagggtattgt caatgaggca gacagattc cgatttgacg ggcaaccaat caatgaaaca 300  
 gacacacctg cacagttgga aatggaggat gaagatacaa ttgatgtgtt ccaacagcag 360  
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<210> 4  
 <211> 95  
 <212> PRT  
 <213> Homo sapiens

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 Lys Ile Lys Arg His Thr Pro Leu Ser Lys Leu Met Lys Ala Tyr Cys  
 35 40 45  
 Glu Arg Gln Gly Leu Ser Met Arg Gln Ile Arg Phe Arg Phe Asp Gly  
 50 55 60  
 Gln Pro Ile Asn Glu Thr Asp Thr Pro Ala Gln Leu Glu Met Glu Asp  
 65 70 75 80  
 Glu Asp Thr Ile Asp Val Phe Gln Gln Gln Thr Gly Gly Val Tyr  
 85 90 95

<210> 5  
 <211> 1733  
 <212> DNA  
 <213> Homo sapiens  
 <220>  
 <221> modified\_base  
 <222> (19)  
 <223> N = A, C, G or T

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 tgtgaagaca gagaatgacc acatcaacct gaaggtggcc gggcaggacg gctccgtggt 180  
 gcagttcaag atcaagaggc acacgtcgct gagcaagctg atgaaggcct actgcgagag 240  
 gcagggttg tcaatgaggc agatcagatt caggttcgac gggcagccaa tcaatgaaac 300  
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 gcccgggccc tccatcctcg cattgctgtt gaatggtgag cacgtgacca tgccgaccac 480  
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 <212> PRT  
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 20 25 30  
 Ile Lys Arg His Thr Ser Leu Ser Lys Leu Met Lys Ala Tyr Cys Glu  
 35 40 45  
 Arg Gln Gly Leu Ser Met Arg Gln Ile Arg Phe Arg Phe Asp Gly Gln  
 50 55 60  
 Pro Ile Asn Glu Thr Asp Thr Pro Ala Gln Leu Arg Met Glu Asp Glu  
 65 70 75 80  
 Asp Thr Ile Asp Val Phe Gln Gln Gln Thr Gly Gly Val Pro Glu Ser  
 85 90 95  
 Ser Leu Ala Gly His Ser Phe  
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<210> 7  
 <211> 9  
 <212> PRT  
 <213> Artificial Sequence  
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 1 5

<210> 8  
 <211> 30  
 <212> DNA  
 <213> Homo sapiens

<400> 8  
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<210> 9

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<211> 30  
<212> DNA  
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<210> 10  
<211> 7  
<212> PRT  
<213> Artificial Sequence  
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Peptide

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<210> 11  
<211> 9  
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<213> Influenza virus

<400> 11  
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1 5

<<210> 12  
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<212> PRT  
<213> Homo sapiens

<400> 12  
His Ser Thr Val  
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<210> 13  
<211> 101  
<212> PRT  
<213> Saccharomyces cerevisiae

<400> 13  
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Glu Val Lys Pro Glu Thr His Ile Asn Leu Lys Val Ser Asp Gly Ser  
20 25 30  
Ser Glu Ile Phe Phe Lys Ile Lys Lys Thr Thr Pro Leu Arg Arg Leu  
35 40 45  
Met Glu Ala Phe Ala Lys Arg Gln Gly Lys Glu Met Asp Ser Leu Arg  
50 55 60  
Phe Leu Tyr Asp Gly Ile Arg Ile Gln Ala Asp Gln Thr Pro Glu Asp  
65 70 75 80  
Leu Asp Met Glu Asp Asn Asp Ile Ile Glu Ala His Arg Glu Gln Ile  
85 90 95  
Gly Gly Ala Thr Tyr

[illegible]

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<400> 15
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Ile Glu Pro Thr Asp Lys Val Glu Arg Ile Lys Glu Arg Val Glu Glu
      20          25          30
Lys Glu Gly Ile Pro Pro Gln Gln Arg Leu Ile Tyr Ser Gly Lys
      35          40          45
Gln Met Asn Asp Glu Lys Thr Ala Ala Asp Tyr Lys Ile Leu Gly Gly
      50          55          60
Ser Val Leu His Leu Val Leu Ala Leu Arg Gly Gly
      65          70          75

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<400> 16
Val Gln Asp Leu Ala Gln Leu Val Glu Ala Thr Gly Val Pro Leu
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Pro Phe Gln Lys Leu Ile Phe Lys Gly Lys Ser Leu Lys Glu
      20      25      30

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